

SEQUENCE LISTING

5 <110> Gure, Ali  
Stockert, Elisabeth  
Scanlan, Matthew J.  
Jager, Dirk  
Old, Lloyd J.  
Chen, Yao-Tseng

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20	gtttttccac	gggatttttt	taaaaaggga	ccncttccc	ngggcntttt	ttttanaaag	1080
	gacccattcc	aantttttgn	ttgnaaaggg	accnttccct	ngggtttnt	aaanngggac	1140
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 <211> 748  
 <212> DNA  
 <213> Homo sapiens

<400> 16

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35	tccacggctt	cgccaacggg	gacgcccgtc	tcagcttcgc	caacggggac	gccgcccagg	300
	gcatcgccaa	cggggacgcc	accaagggca	tgggcaacga	ggtcaccatc	cacggcatcg	360
	ctaacgagga	cgccgtccag	ggcatcgcta	acgaggtggc	cgcccagggc	atcgccaacg	420
	aggacgccgc	ccagggaatc	gccgaggatg	tcgcacaggg	catcgccaac	gaggacgccg	480
	cccagggcat	cgccaacaag	gaggccgccc	agggcatcgc	caacgaggac	gccgcccagg	540
40	gaatcgctga	ggacgtcgca	cagggcatcg	ccaacgagga	tgccgcccag	ggcatcgcca	600
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<210> 17  
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<400> 17

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55	ctnaggctga	ggttggggct	ggggctggng	ctgacgctgg	ggctgaggct	nggnctgagg	300
	ctggagctgg	ggctgangct	ggggctgggg	ctgnngctga	nctggggctg	aggctccngc	360
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	tgnngnatgn	ctgnngctnn	cntccnngac	aaananttnn	aacttngngt	ttnttcctgg	480
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15 <400> 18

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 35 40 45  
 Asp Arg Val Lys Arg Pro Met Asn Ala Phe Met Val Trp Ser Arg Gly  
 50 55 60  
 Gln Arg Arg Lys Met Ala Gln Glu Asn Pro Lys Met His Asn Ser Glu  
 65 70 75 80  
 Ile Ser Lys Arg Leu Gly Ala Glu Trp Lys Val Met Ser Glu Ala Glu  
 85 90 95  
 Lys Arg Pro Phe Ile Asp Glu Ala Lys Arg Leu Arg Ala Leu His Met  
 100 105 110  
 30 Lys Glu His Pro Asp Tyr Lys Tyr Arg Pro Arg Arg Lys Thr Lys Thr  
 115 120 125  
 Leu Leu Lys Lys Asp Lys Tyr Ser Leu Ala Gly Gly Leu Leu Ala Ala  
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 Gly Ala Gly Gly Gly Gly Ala Ala Val Ala Met Gly Val Gly Val Gly  
 145 150 155 160  
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 Ala Gly Gly Ala Tyr Ala His Val Asn Gly Trp Ala Asn Gly Ala Tyr  
 180 185 190  
 40 Pro Gly Ser Val Ala Ala Ala Ala Ala Ala Ala Met Met Gln Glu  
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 Ala Gln Leu Ala Tyr Gly Gln His Pro Gly Ala Gly Gly Ala His Pro  
 210 215 220  
 His Arg Thr Pro Ala His Pro His Pro His Pro His Ala His Pro  
 225 230 235 240  
 45 His Asn Pro Gln Pro Met His Arg Tyr Asp Met Gly Ala Leu Gln Tyr  
 245 250 255  
 Ser Pro Ile Ser Asn Ser Gln Gly Tyr Met Ser Ala Ser Pro Ser Gly  
 260 265 270  
 50 Tyr Gly Gly Leu Pro Tyr Gly Ala Ala Ala Ala Ala Ala Ala His  
 275 280 285  
 Gln Asn Ser Ala Val Ala Ala Ala Ala Ala Ala Ala Ala Ser Ser  
 290 295 300  
 Gly Ala Leu Gly Ala Leu Gly Ser Leu Val Lys Ser Glu Pro Ser Gly  
 305 310 315 320  
 55 Ser Pro Pro Ala Pro Ala His Ser Arg Ala Pro Cys Pro Gly Asp Leu  
 325 330 335  
 Arg Glu Met Ile Ser Met Tyr Leu Pro Ala Gly Glu Gly Gly Asp Pro  
 340 345 350  
 60 Ala Ala Ala Ala Ala Ala Ala Gln Ser Arg Leu His Ser Leu Pro  
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Thr His Ile  
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5 <210> 19  
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15 35 40 45  
Met Val Trp Ser Arg Gly Gln Arg Arg Lys Met Ala Gln Glu Asn Pro  
50 55 60  
Lys Met His Asn Ser Glu Ile Ser Lys Arg Leu Gly Ala Glu Trp Lys  
65 70 75 80  
20 Leu Leu Ser Glu Thr Glu Lys Arg Pro Phe Ile Asp Glu Ala Lys Arg  
85 90 95  
Leu Arg Ala Leu His Met Lys Glu His Pro Asp Tyr Lys Tyr Arg Pro  
100 105 110  
Arg Arg Lys Thr Lys Thr Leu Met Lys Lys Asp Lys Tyr Thr Leu Pro  
115 120 125  
25 Gly Gly Leu Leu Ala Pro Gly Gly Asn Ser Met Ala Ser Gly Val Gly  
130 135 140  
Val Gly Ala Gly Leu Gly Ala Gly Val Asn Gln Arg Met Asp Ser Tyr  
145 150 155 160  
30 Ala His Met Asn Gly Trp Ser Asn Gly Ser Tyr Ser Met Met Gln Asp  
165 170 175  
Gln Leu Gly Tyr Pro Gln His Pro Gly Leu Asn Ala His Gly Ala Ala  
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Gln Met Gln Pro Met His Arg Tyr Asp Val Ser Ala Leu Gln Tyr Asn  
195 200 205  
35 Ser Met Thr Ser Ser Gln Thr Tyr Met Asn Gly Ser Pro Thr Tyr Ser  
210 215 220  
Met Ser Tyr Ser Gln Gln Gly Thr Pro Gly Met Ala Leu Gly Ser Met  
225 230 235 240  
40 Gly Ser Val Val Lys Ser Glu Ala Ser Ser Ser Pro Pro Val Val Thr  
245 250 255  
Ser Ser Ser His Ser Arg Ala Pro Cys Gln Ala Gly Asp Leu Arg Asp  
260 265 270  
Met Ile Ser Met Tyr Leu Pro Gly Ala Glu Val Pro Glu Pro Ala Ala  
275 280 285  
45 Pro Ser Arg Leu His Met Ser Gln His Tyr Gln Ser Gly Pro Val Pro  
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Gly Thr Ala Ile Asn Gly Thr Leu Pro Leu Ser His Met  
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50 <210> 20  
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<212> PRT  
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Asp Ser Leu Ala His Arg Pro Pro Ser Ser Ala Pro Thr Glu Ser Gln  
35 40 45  
60 Gly Leu Phe Thr Val Ala Ala Pro Ala Pro Gly Ala Pro Ser Pro Pro

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40